SEQUENCE LISTING

<110> Roses, Allen David

<120> Iterative Analysis of Non-Responding
   Populations in the Design of Pharmacogenetic Studies

<130> PU3948US3

<140> to be assigned
<141> concurrently

<150> 60/194,789
<151> 2000-04-05

<160> 2

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 484
<212> DNA
<213> Homo Sapiens

<400> 1
ggcgttgccg ctctgaatgc cagccctaac ccctaatgtc cctactgcag cctcccagca 60
tcccccctgc aacctcccag caactccctg tacccctcct aggatcgctc ctgcatcccc 120
cattatcccc cccttcactc ctcgcggcat cccccctgca cccccagcat cccccctgca 180
gcccttccag catccccctg cacctctccc aggatctccc ctgcaacccc cattatcccc 240
cctgcacccc tcgcagtatc ccccctgcac ccccagcat cccccatgc accccggca 300
tccccctgc acccctccag cattctcctt gcaccctacc agtattcccc cgcatcccgg 360
cctccaagcc tcccgcccac cttgcggtcc ccgccctggc gtctaggtgg caccagaatc 420
ccgcgcggac tccacccgct gggagctgcc ctcgcttgcc cgtggttgtc cagctcagtc 480
cctc   484

<210> 2
<211> 528
<212> DNA
<213> Homo Sapiens

<400> 2
ggcgttgccg ctctgaatgc cagccctaac ccctaatgtc cctactgcag cctcccagca 60
tcccccctgc aacctcccag caactccctg tacccctcct aggatcgctc ctgcatcccc 120
cattatcccc cccttcactc ctcgcggcat cccccctgca cccccagca tcccccctgc 180
agcccccccca gcatctcccc tgcaccccca gcatccccc tgcagcccctt ccagcatccc 240
cctgcacctc tccaggatc tccctgcaa cccccattat cccccctgca ccctcgcag 300
tatcccccct gcaccccca gcatcccccc atgcacccc ggcatccccc ctgcacccct 360
ccagcattct ccttgcaccc taccagtatt ccccgcatc ccggcctcca agcctcccgc 420
ccaccttgcg gtccccgccc tggcgtctag gtggcaccag aatcccgcgc ggactccacc 480
cgctgggagc tgccctcgct tgcccgtggt tgtccagctc agtccctc   528